

In the Claims:

1. (Currently amended) A method of generating, storing and accessing genomic information, that comprises, generating image patterns containing such genomic information; standardizing the format of the image patterns as standardized two-dimensional spot image patterns, and storing the standardized two-dimensional image spot patterns in a database library; and accessing the database for such purposes as inputting additional of such standardized format image patterns for storage in the database to develop the same, retrieving specified image pattern information stored in the database, and image pattern comparison and analysis amongst said standardized format image patterns.
2. (Currently amended) A method of generating, storing and accessing genomic information, that comprises, generating image patterns containing such information in the form of standardized gene-specific two-dimensional spot patterns of gene fragments distributed by electrophoresis in a gel; formatting the image spot patterns for standardized entry into an image database; storing the same in said database; and enabling accessing of the database for such purposes as the inputting of new image standardized two-dimensional format spot patterns into the database, the retrieving of specific standardized image spot pattern information stored in the data base, and comparison and analysis amongst the standardized two-dimensional format image spot patterns.
3. (Original) The method of claim 2 wherein the electrophoretic distribution in the gel is effected, following multiplex PCR operation on the gene fragments, by two-dimensional gene scanning electrophoresis.
4. (Original) The method of claim 2 wherein one or more of the inputting, retrieving and accessing is effected by external communication with data base constructors and users.
5. (Original) The method of claim 4 wherein said communication is over the internet.
6. (Currently amended) The method of claim 2 wherein the electrophoresis is enabled by providing gene-specific customized assay kits to the electrophoresis user, designed to facilitate said standardized formatting of the resulting uniform two-dimensional spot pattern images.

7. (Original) The method of claim 6 wherein each kit is customized for a specific known gene or genes of specific individuals.
8. (Original) The method of claim 6 wherein the kit is designed for an unknown gene or genes.
9. (Original) The method of claim 7, wherein the image comparison and analyzing assists in identifying known-mutations of the specified gene(s).
10. (Original) The method of claim 7 wherein spot image comparison and analyzing assists in identifying target populations for new drugs candidacy.
11. (Original) The method of claim 6 wherein the inputting of new spot pattern images into the database, and the retrieving and accessing of information therefrom is networked to external researchers, diagnosticians and others, enabling the developing of the data base from global population bases, and usage of the data base globally as well.
12. (Currently amended) A system for generating, storing and accessing genomic information, having in combination, apparatus for generating image patterns containing such information in the format of gene-specific uniform two-dimensional spot patterns of gene fragments distributed by electrophoresis on a gel; software designed for formatting the fragment spot patterns for standardized entry into an image database; a database storage and retrieval apparatus for storing the spot pattern images stored in the database; and means for enabling remote accessing of the database for such purposes as the inputting of new standardized two-dimensional format image spot patterns into the database, the remote retrieving of specific spot pattern information stored in the database, and the comparison and analysis amongst the standardized two-dimensional format image spot patterns.
13. (Original) The system of claim 12 wherein the electrophoretic distribution in the gel is effected, following multiplex PCR on the gene fragments, by two-dimensional gene scanning electrophoresis apparatus.
14. (Original) The system of claim 12 wherein one or more of the inputting, retrieving and accessing is effected by an external two-way communication network provided for and with data base constructors and users.

15. (Original) The system of claim 14 wherein said communication network is over the internet.

16. (Currently amended) A new method of globally doing business in the sale of services and products related to genomic information, that comprises, creating a database library of standardized two-dimensional uniform formatted electrophoresis gel spot pattern images of specified gene fragments derived from customer contributors to, and users of, the database library; providing customized assay kit products to such customers designed to insure standardization of such formatted spot pattern images for storage in the database library; enabling two-way communication preferably over the internet, between customers and users and the database library for enabling (1) customer or user remote inputting of new standardized two-dimensional format electrophoretic spot pattern images of specified genes and individuals, resulting from population studies and/or from diagnostic testing research; (2) providing for communication to such customers and users, comparison and analyzing services of inputted standardized format spot pattern images relative to images stored in the data base; and (3) providing spot pattern image information stored in the database, on request, to such customers and users.

17. (Original) The method of claim 2 in which the compiled 2-D spot pattern images are directly correlated with protein structure through the linking of the database to protein databases and protein modeling software.

18. (Original) The method of claim 1 wherein the database is linked to other bioinformatics resources including other genomic references and protein modeling software and databases.

19. (Original) The method of claim 2 wherein the database is linked to other bioinformatics resources including other genomic references and protein modeling software and databases.

20. (Original) The method of claim 16 wherein the database is linked to other bioinformatic resources including other genomic references and protein modeling software and databases.

21. (Previously added) The method of claim 16 wherein the standardized format of the electrophoresis gel spot pattern images is formed by standardizing size and contrast.

22. (Previously added) The method of claim 1 wherein the standardized format of the image patterns is formed by standardizing size and contrast.